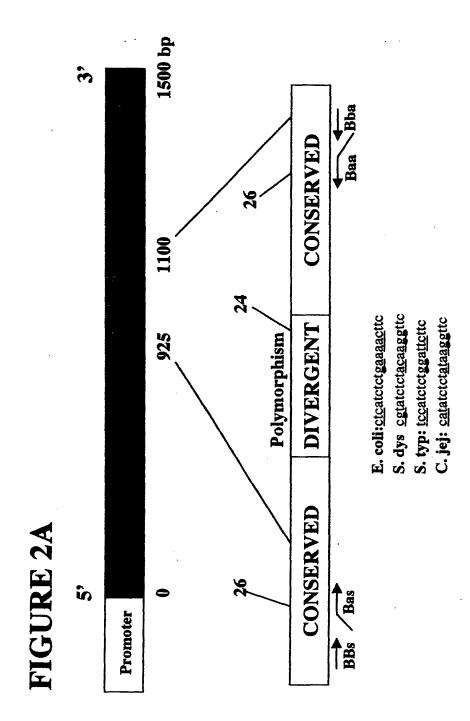


FIG. 1B

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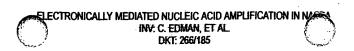
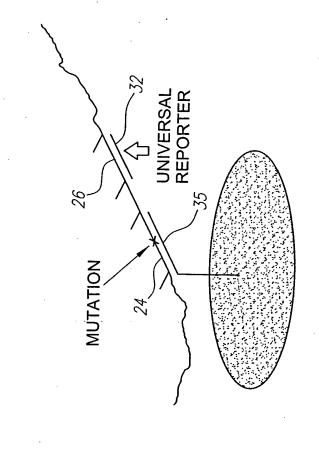


FIGURE 2B

Control E.coli 0157:H7 Salmonella Shigella M1 M2



SEQ SPECIFIC REPORTER

-28

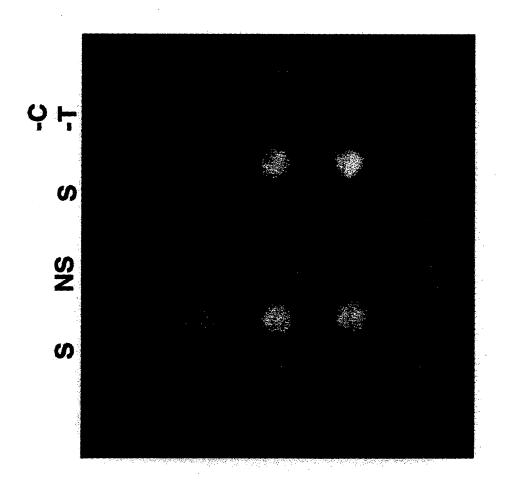
MUTATION 24

SEQUENCE-SPECIFIC CAPTURE

UNIVERSAL CAPTURE

FIG. 2C

FIG. 2D



Salmonella

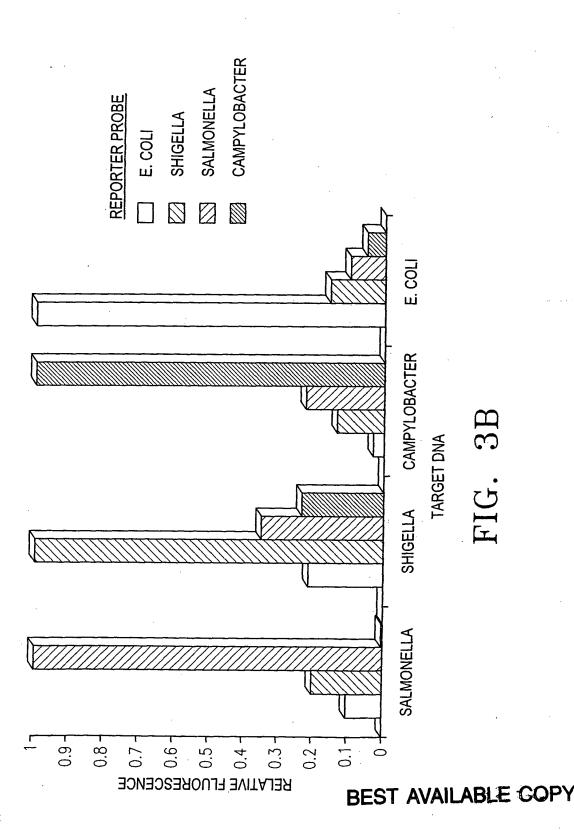
Salmonella

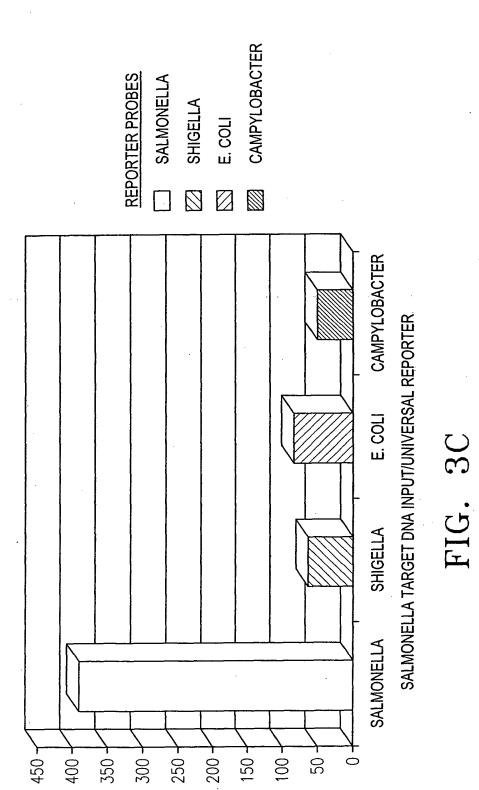
Campylobacter

FIGURE 3A

Shigella







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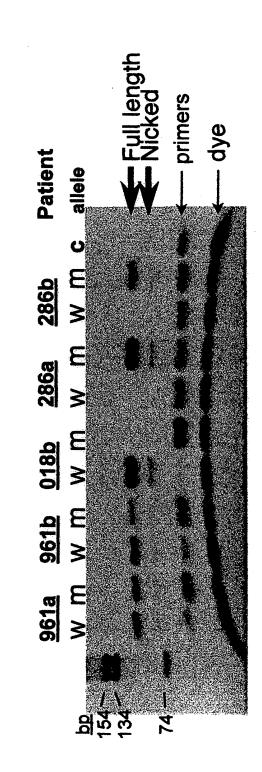
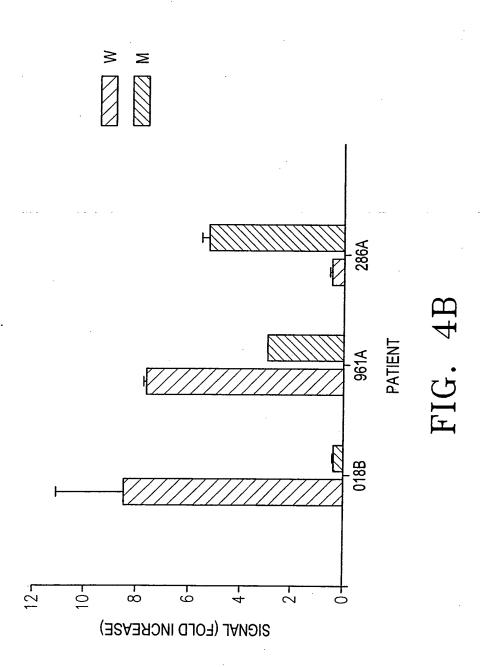
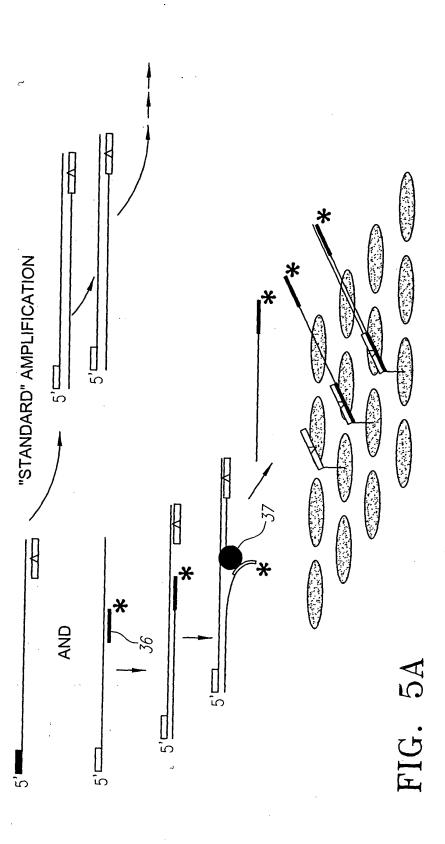


FIGURE 4A



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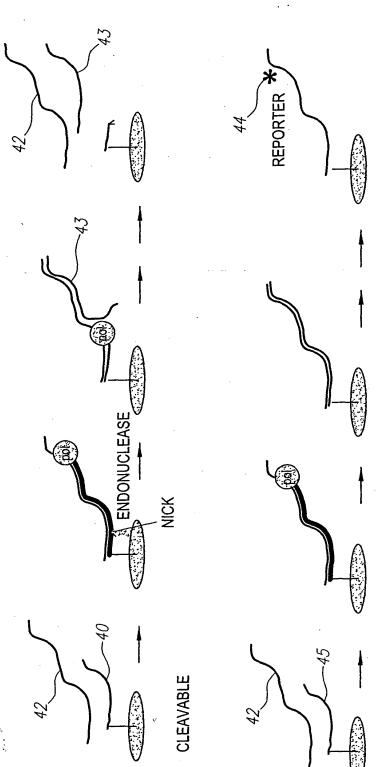
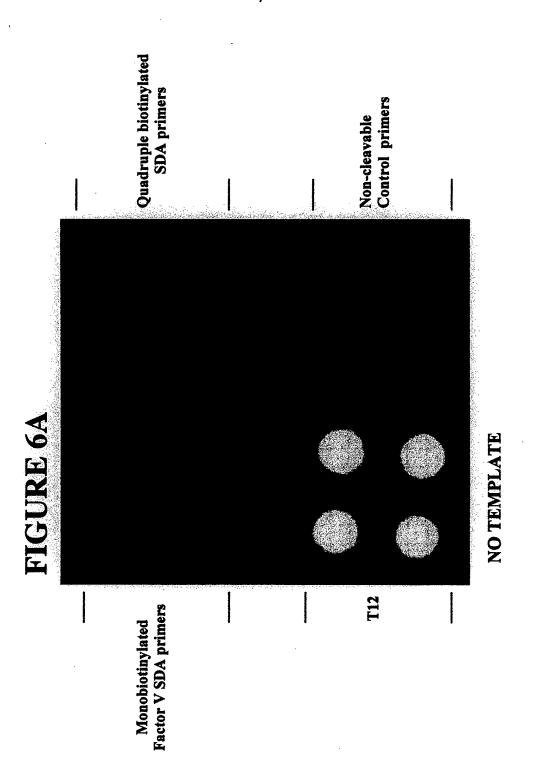


FIG. 5B

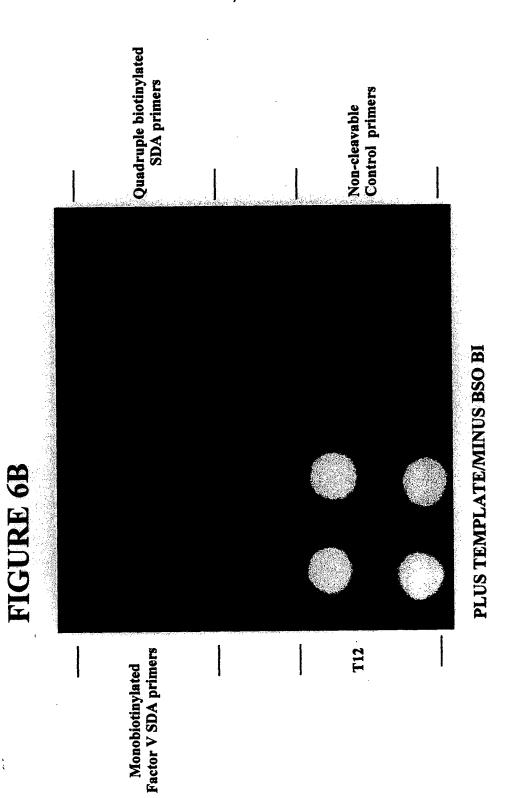
NONCLEAVABLE

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FLECTRONICALLY MEDIATED NUCLEIC ACID AMPLIFICATION IN NACRA INV: C. EDMAN, ET AL DKT: 266/185

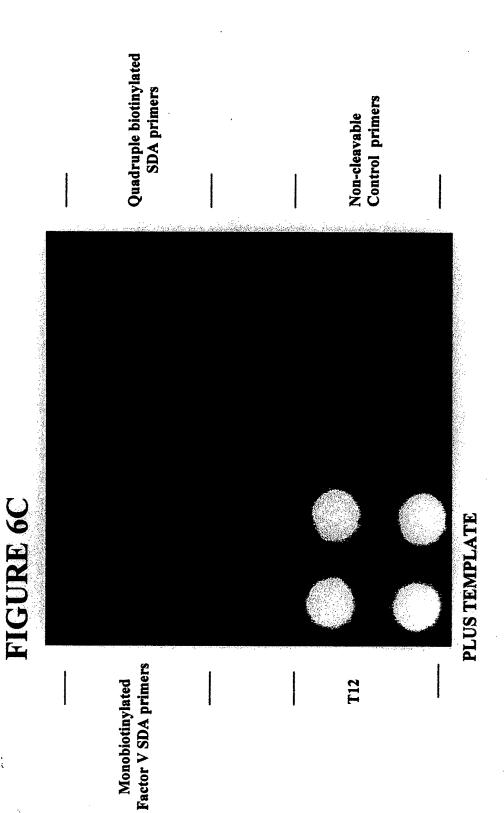
13/34

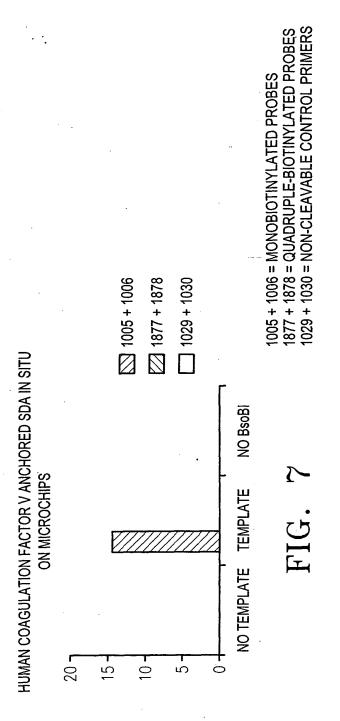


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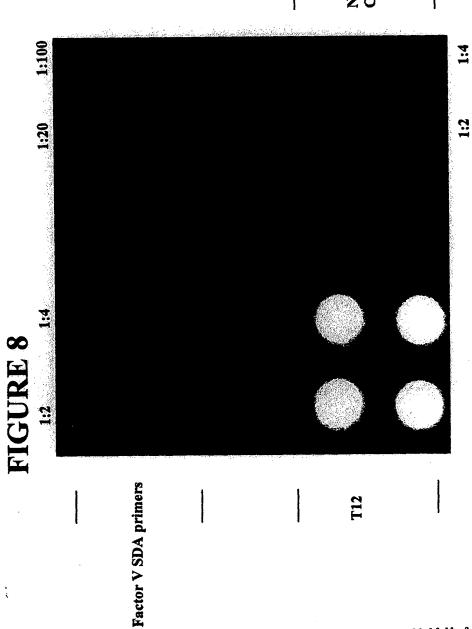
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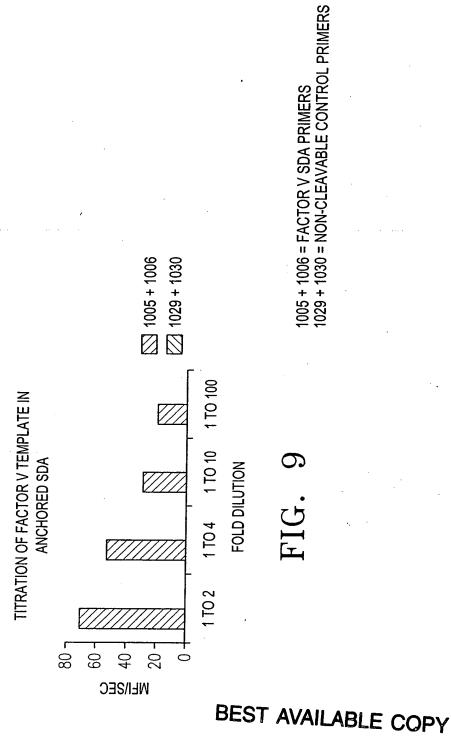
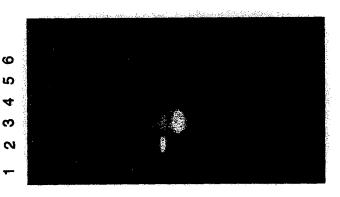


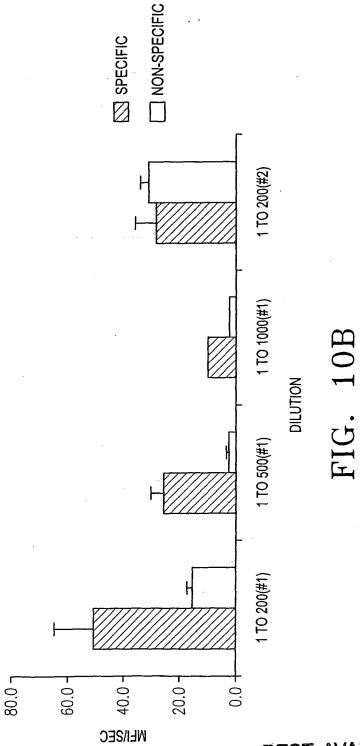
FIGURE 10A

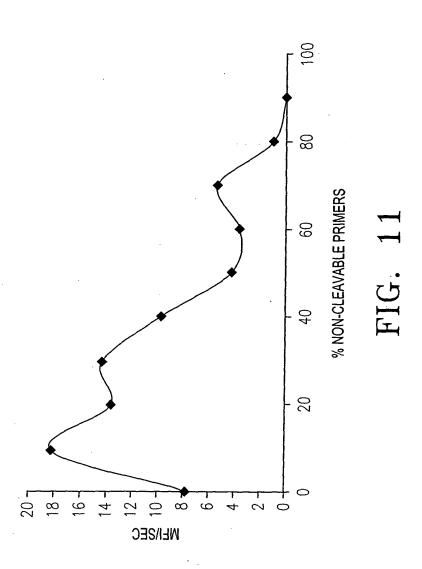
Lane

- 1. mol wt markers
- 2. PCR amplification 3. NASBA 1X templa
- 3. NASBA 1X template
- NASBA 1,000X diluted template
 NASBA 1,000,000X diluted template

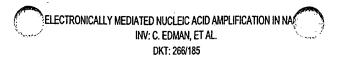
6. NASBA - no template







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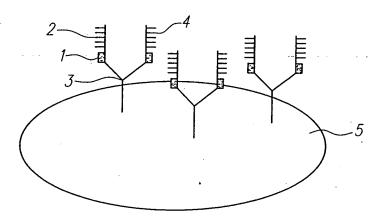
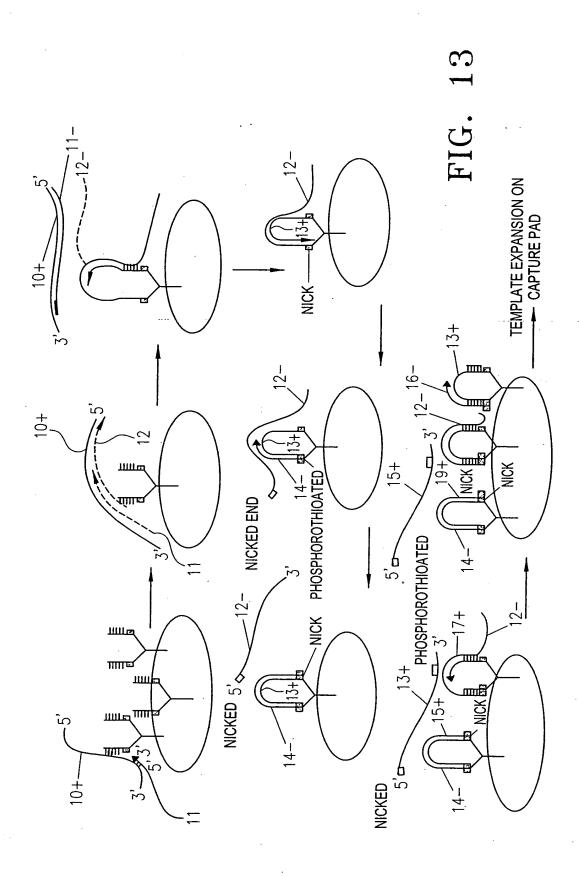
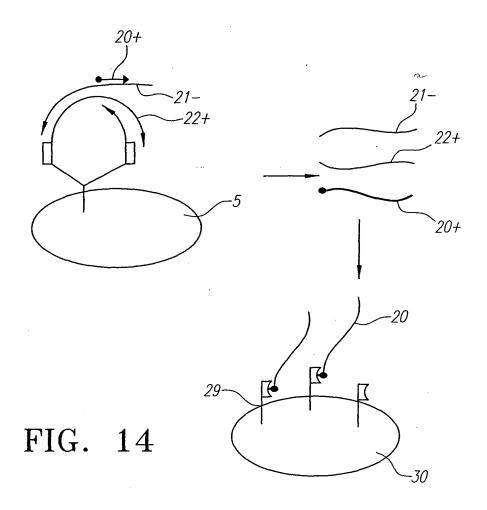


FIG. 12







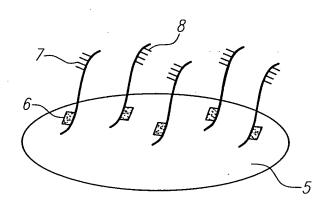


FIG. 15



FIGURE 16

FIGURE 17 E CHI HC HC

Control - No template + all reporter oligos

Experimental Layout

ATA

ATA

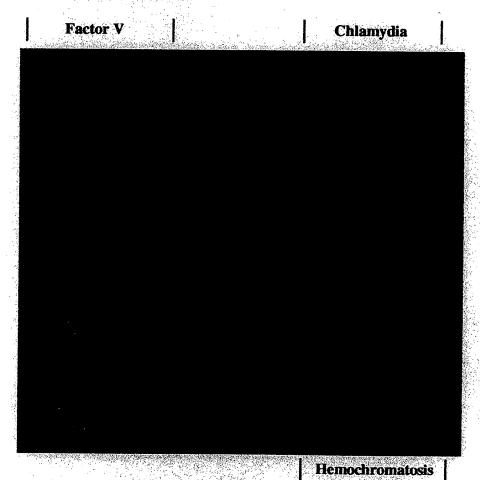
BEST AVAILABLE COPY

All templates + Factor V Reporter oligo

All templates + Factor V, Chlamydia

Reporter Oligos

FIGURE 20



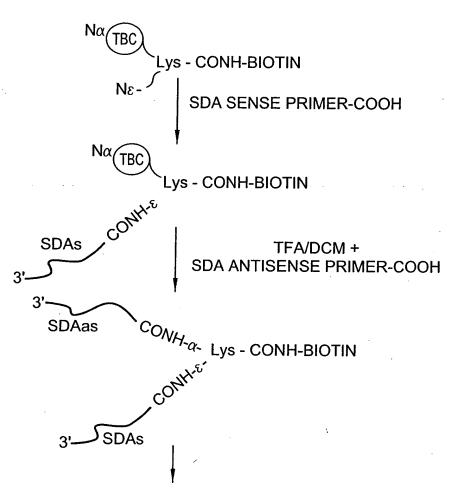
All templates + Factor V, Chlamydia and Hemachromatosis reporter oligos

FIGURE 21

Factor V Chlamydia Hemochrom ALL

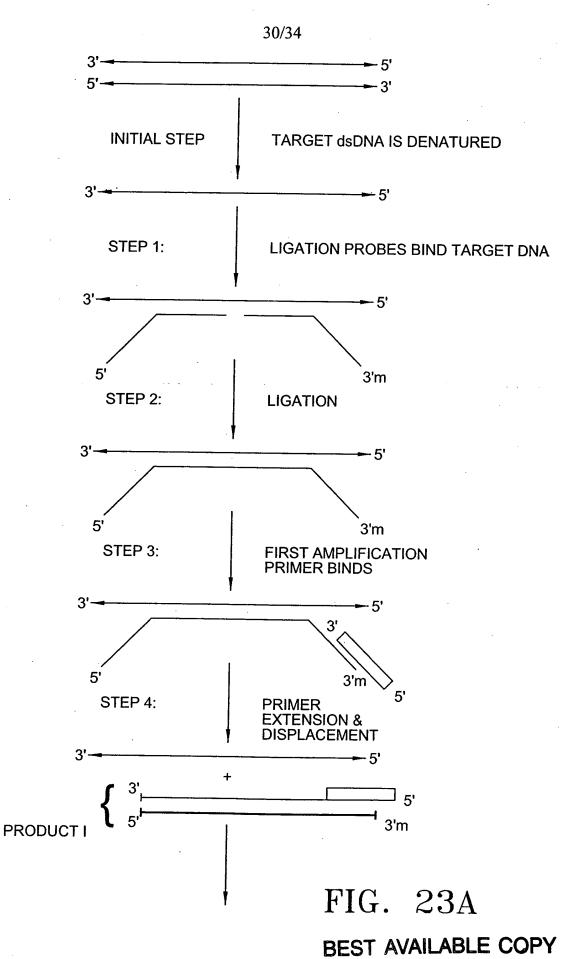
- + - + - + - + - +

Control Solution SDA reactions



ATTACH TO STREPTAVIDIN PERMEATION LAYER ON MICROCHIP

FIG. 22



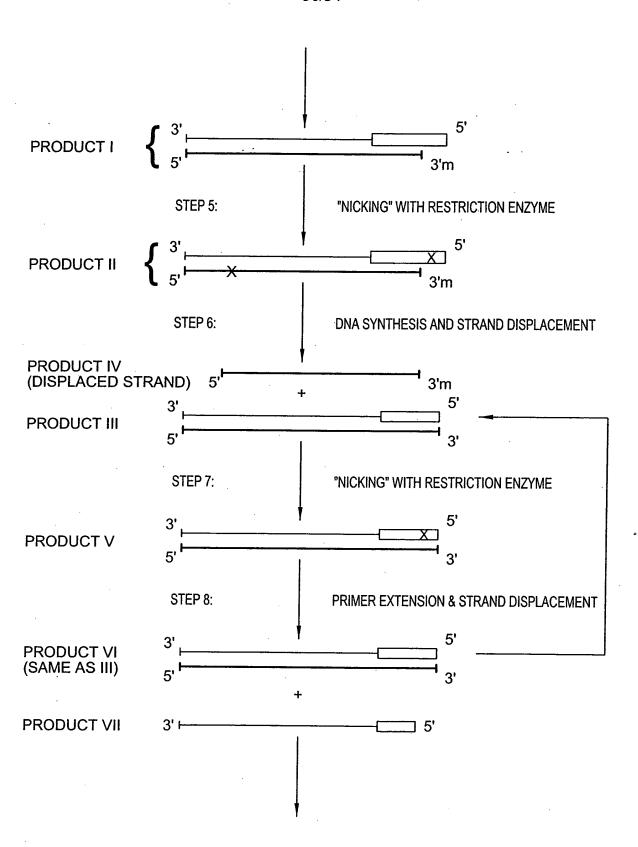
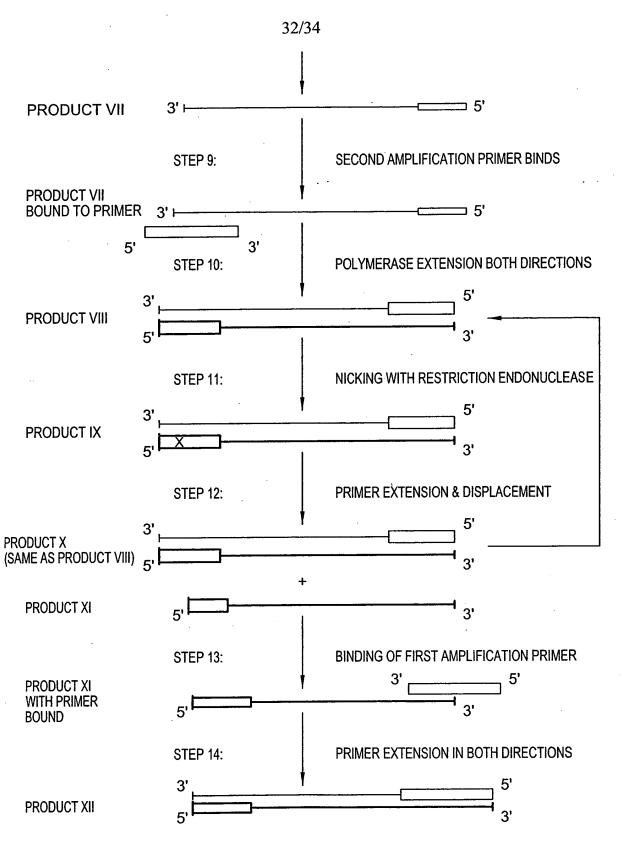


FIG. 23B





(PRODUCT XII CAN RE-ENTER PATHWAY AND BE FURTHER AMPLIFIED IN A MANNER SIMILAR TO PRODUCT III, FOLLOWING STEP 6)

FIG. 23C BEST AVAILABLE COPY

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LIGATION-DEPENDENT DETECTION OF THE SALMONELLA spaQ GENE

IGATION PROBES LP1 AND LP2:

5-nnnnncaacatgacatcattacgagacgggatagttaaatggatgatgatttagtgnnnn-3' spaQ1 TEMPLATE $LP1^2$

3-*aattccgcatgagctgggtaatgttgtactgtagtaatgctctgc*-5

3'-cctatcaatttacctactaaatcacgattatcccctagagtcatgtgggctc

LP23

ttcagacctcgccttagc-5'

MPLIFICATION PRIMER SEQUENCES S1 AND S2:

3-*aattccgcatgagctgggtaatgttgtactgtagtaatgctctgc*-5'

댎

5'-accgcatcgaatgcatgtctcgggtaaggcgtactcgacc

S14

3'-cetateaatttacetactaaateacgattateeectagagteatgtgggetetteagaeetegeettage-5' LP2

 -52^{5}



